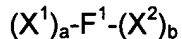


Claims

1. (Previously amended) A composition of matter of the formula



and multimers thereof, wherein:

F^1 is an Fc domain;

X^1 and X^2 are each independently selected from $-(L^1)_c-P^1$, $-(L^1)_c-P^1-(L^2)_d-P^2$, $-(L^1)_c-P^1-(L^2)_d-P^2-(L^3)_e-P^3$, and $-(L^1)_c-P^1-(L^2)_d-P^2-(L^3)_e-P^3-(L^4)_f-P^4$

P^1 , P^2 , P^3 , and P^4 are each independently randomized Ang-2 binding peptide sequences;

L^1 , L^2 , L^3 , and L^4 are each independently linkers; and

a , b , c , d , e , and f are each independently 0 or 1, provided that at least one of a and b is 1; and

wherein "peptide" refers to molecules of 2 to 40 amino acids and wherein neither X^1 nor X^2 is a native protein.

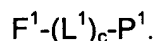
2. (Original) The composition of matter of Claim 1 of the formulae



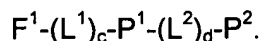
or



3. (Original) The composition of matter of Claim 1 of the formula



4. (Original) The composition of matter of Claim 1 of the formula



5. (Original) The composition of matter of Claim 1 wherein F^1 is an IgG Fc domain.

6. (Original) The composition of matter of Claim 1 wherein F^1 is an IgG1 Fc domain.

7. (Original) The composition of matter of Claim 1 wherein F^1 comprises the sequence of SEQ ID NO: 2.

8-62. (Previously canceled)

63. (Previously presented) The composition of matter of Claim 3, wherein P¹ is selected by phage display, *E. coli* display, ribosome display, RNA-peptide screening, yeast-based screening, or chemical-peptide screening.

64. (Previously presented) The composition of matter of Claim 4, wherein P¹ and P² are selected by phage display, *E. coli* display, ribosome display, RNA-peptide screening, yeast-based screening, or chemical-peptide screening.